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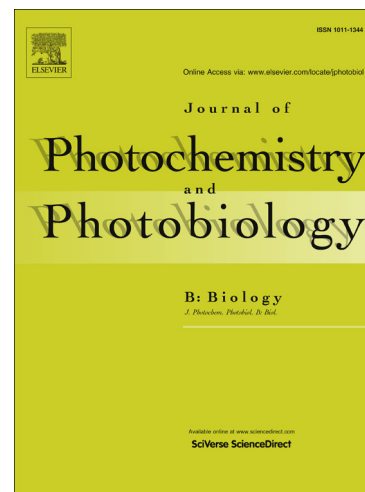
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In vivo estimation of pigment composition and optical absorption cross-section by spectroradiometry in four aquatic photosynthetic micro-organisms

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Abstract

The objective of the present study was to estimate *in vivo* pigment composition and to retrieve absorption cross-section values, a^* , of photosynthetic micro-organisms using a non-invasive technique of reflectance spectrometry. To test the methodology, organisms from different taxonomical groups and different pigment composition were used (*Spirulina platensis* a Cyanophyta, *Porphyridium cruentum* a Rhodophyta, *Dunaliella tertiolecta* a Chlorophyta and *Entomoneis paludosa* a Bacillariophyta) and photoacclimated to two different irradiance levels: 25 $\mu\text{mol photon.m}^{-2}.\text{s}^{-1}$ (Low Light, LL) and 500 $\mu\text{mol photon.m}^{-2}.\text{s}^{-1}$ (High Light, HL). Second derivative spectra from reflectance were used to identify pigment *in vivo* absorption bands that were linked to specific pigments detected by High Performance Liquid Chromatography. Whereas some absorption bands such as those induced by Chlorophyll (Chl) *a* (416, 440, 625 and around 675 nm) were ubiquitous, others were taxonomically specific (e.g. 636 nm for Chl *c* in *E. paludosa*) and/or photo-physiological dependent (e.g. 489 nm for zeaxanthin in the HL-acclimated *S. platensis*). The optical absorption cross-section, a^* , was retrieved from reflectance data using a radiative transfer model previously developed for microphytobenthos. Despite the cellular Chl *a* decrease observed from LL to HL (up to 88 % for *S. platensis*), the a^* increased, except for *P. cruentum*. This was attributed to a 'package effect' and to a greater absorption by photoprotective carotenoids that did not contribute to the energy transfer to the core Chl *a*.

Key-words: microalgae, cyanobacteria, *in vivo* absorption bands, reflectance, pigments, absorption cross-section.

1 Introduction

Marine coastal areas, particularly bays and estuaries, are amongst the most productive ecosystems on Earth (e.g. [1]). Polyphyletic photosynthetic microbial

communities (micro-algae, phytoplankton and microphytobenthos, and cyanobacteria) support this high coastal productivity. Their ecological success and adaptability to light variations in such ecosystems are partially explained by the wide plasticity of their photosynthetic apparatus constituted by a variety of light-harvesting proteins and pigments. This variability reflects the large range of photoecological responses of these micro-organisms due to different photoadaptation characteristics and photoacclimation processes [2]. An important consequence is a variability in primary production rates, depending of species and their respective ecophysiology. Estimation of *in vivo* cell absorption properties in the photosynthetic active radiation (PAR) domain (400 to 700 nm) provides important taxonomic and ecophysiological information. Namely, pigment composition allows to assess the community structure and physiology, whereas the optical absorption cross-section, i.e. a^* , provides information about the potential primary production [3]. Frequently, this information is retrieved from absorption signatures measured either *in vitro* (i.e. in solvent-extracted pigments) or *in vivo* (i.e. in intact cells) [3]. Pigment absorption properties are dependent on their molecular environment thus the *in vitro* estimations are subject to important constraints [4, 5]. Firstly, pigment absorption maxima usually shifts from *in vitro* measurements in comparison to *in vivo* measurements. This shift can vary from a few to hundreds of nanometers, depending on the pigment, the association with protein complexes (i.e. Light Harvesting Complexes, LHC) and the pigment location inside the cell [4-8]. Consequently, the assignment of each absorption band to its respective pigment is very difficult to achieve [4]. Secondly, *in vivo* absorption is always lower than *in vitro* leading to a misestimation of a^* [2]. To overcome these constraints, *in vivo* measurements of absorbance signatures can be performed by spectrophotometry, either on cell suspensions or on thylakoids and isolated LHCs [9,

10]. Spectrophotometry assumes that the sample is homogenous, and that cell concentration (or thylakoids or LHCs) is high enough to detect absorption, but low enough to avoid self-shading. Furthermore, the absorption measurements can be affected by light scattering caused by cells in suspension [3]. Some improvements have been proposed, including the use of an integrative sphere or placing the sample (cells on wet glass-fiber filters) close to the detector [5, 11-13]. However, these methods are constrained by the need for β -correction algorithms to remove the path length amplification effect due to the filter [12]. Therefore, there is a growing interest in using non-destructive techniques to investigate micro-algal absorption properties while simultaneously limiting the scattering effect. Recently, reflectance spectroradiometric techniques have been used to assess micro-algal pigment composition [7, 14-17]. Reflectance spectra can be studied using derivative analysis, a technique which is particularly useful in removing the scattering effect from absorption spectra and in resolving overlapping spectral features [18]. Derivative analysis have been used to detect absorption bands due to electronic transitions [9] and to assess the light-dependent (de-)epoxidation of xanthophylls [10, 16]. Despite the great potential of this technique, few studies have used it to assess the pigment content of micro-algae and to investigate all pigment specific absorption bands [16, 19-21]. Recently, Kazemipour et al. [14] developed a radiative transfer model (MPBOM, MicroPhytoBenthos Optical Model) which allows the determination of microphytobenthos biofilm absorption coefficients using reflectance measurements without destroying the biofilm. This dimensionless coefficient corresponds to the optical cross-section, a^* , when it is divided by the chlorophyll (Chl) a concentration ($\text{m}^2.\text{mg Chl}a^{-1}$). Initially developed for benthic microalgae, this model can be applied to any other photosynthetic microorganisms.

The objective of the present study was to use reflectance data to estimate *in vivo* pigment composition and determination of a^* values from different photosynthetic microbial species. Pigment composition was estimated from spectral reflectance using second derivative analysis and a^* was calculated using the MPBOM radiative transfer model. To validate the methods, a variety of microbial species were investigated: one Cyanophyta, one Rhodophyta, one Chlorophyta and one Bacillariophyta (diatom). Intraspecific variability was also explored by cultivating all species at two different irradiance levels expected to induce photoacclimation pigment changes at the cell level.

2 Materials and methods

2.1 Biological material

Three species of eukaryotic micro-algae and one of cyanobacteria were chosen for their different pigment signatures: one 'green' species, *Dunaliella tertiolecta* (Chlorophyceae, Chlorophyta), one 'brown' species, *Entomoneis paludosa* (Diatoms, Heterokontophyta), one 'blue', *Spirulina platensis* (Oscillatoriaceae, Cyanophyta) and one 'red' species, *Porphyridium cruentum* (Bangiophyceae, Rhodophyta). *E. paludosa* and *P. cruentum* were obtained from the Nantes Culture Collection (NCC18.2 and NCC48, respectively), *D. tertiolecta* was obtained from the Provasoli-Guillard National Center for Culture of Marine Phytoplankton (CCMP364) and *S. platensis* was obtained from the Alpha Biotech Company (SARL, Asserac 44410, France).

E. paludosa, *P. cruentum* and *D. tertiolecta* species were grown in f/2 medium [22]. *S. platensis* was grown in Zarrouk medium [23]. Three replicates of each species were

grown in a temperature-controlled culture room ($14 \pm 1^\circ\text{C}$, 14/10 h light/dark cycle) in 500-mL glass flasks containing 250 mL of medium at two PAR levels: $25 \mu\text{mol photon.m}^{-2}.\text{s}^{-1}$ (Low Light, LL) and $500 \mu\text{mol photon.m}^{-2}.\text{s}^{-1}$ (High Light, HL). Each culture was acclimated during 10 days, and regular dilution of the batch culture with fresh medium allowed to keep cells into exponential stage and minimize self-shading. A thin compact layer of micro-algal cells, hereafter named 'experimental biofilm', was deposited on appropriate filters according to Kazemipour et al. [14]. For each culture, 5 mL were gently filtered (<1 bar) through glass microfiber filters (GF/C, 25 mm Ø, $1.2 \mu\text{m}$ porosity). This volume and the corresponding cell concentrations were sufficient to form a thin and homogenous compact layer of cells covering the filter surface ($\sim 1.8 \text{ cm}^2$) while reducing the distance between cells and limiting light scattering [14]. To estimate the amount of cells on filters, each culture was sampled. Before any reflectance measurement was carried out, the experimental biofilms were placed 10 min under the corresponding PAR levels (i.e. HL or LL) to minimize rapid changes in pigment composition due to stress. For each PAR level, two filters were used for each culture run in triplicate ($n=6$). Biofilm desiccation during measurements was minimized by maintaining the filters on absorbent paper soaked with culture medium. Reflectance spectra were standardized with reference filters obtained by filtering 5 mL of the corresponding medium alone.

2.2 Radiometric measurements and interpretations

Radiometric measurements were performed using an ASD Fieldspec3 FR[®] portable spectroradiometer that measures radiance ($\text{mW.cm}^{-2}.\text{nm}^{-2}.\text{sr}^{-1}$) in the visible-infrared region of the light spectrum (between 400 and 2500 nm), with a 3 nm spectral resolution and a 1.4 nm sampling interval in the 400-1000 nm wavelength range. The

light source was provided by an internal halogen lamp and the distance between the biofilm and the ASD optical fiber was kept constant by using the ASD High Intensity Contact Probe. This procedure minimizes errors associated with stray light. Reflectance (ρ , dimensionless) was calculated as the ratio between the radiance of the cells on filters and the incident radiance measured on a perfect diffuser (Spectralon® 99%). Ten spectra were recorded and averaged for each filter. Reflectance was standardized (ρ_{std}) to facilitate comparisons between the spectral signatures of cells acclimated to different light conditions following Eq. (1):

$$\rho_{std} = \frac{\rho}{\rho_F} \times \frac{1}{\rho_{925}} \quad (1)$$

where ρ_{std} is the standardized reflectance, ρ is the reflectance of the experimental biofilm, ρ_F is the reflectance of a wet filter with no biofilm, and ρ_{925} is the reflectance of the experimental biofilm at 925 nm which is insensitive to the pigment content of cells [7, 14].

Second derivative spectra ($\delta\delta$) were calculated following Jesus et al. [16, 19] and second derivative peaks were used to assign pigments to absorption wavelengths. To facilitate the comparison between species and light-acclimation conditions, the second derivative values were standardized ($\delta\delta_{std}$), following Eq. (2), to the maximum value between 650 and 690 ($\delta\delta_{red}$), which is characteristic of Chl *a* red absorption:

$$\delta\delta_{std} = \frac{\delta\delta}{\delta\delta_{red}} \quad (2)$$

2.3 Pigment analysis by High Performance Liquid Chromatography (HPLC)

Following radiometric measurements, the filters were immediately frozen in liquid nitrogen, and stored at -80 °C until pigment extraction and analysis. Pigment analysis

was performed on 3 filters for each culture and each PAR level ($n=3$) with a Waters® HPLC device as described in Méléder et al. [7]. Pigments were extracted after thawing in 90% acetone, ground and left for 24 h in the dark at +4°C as recommended by Mantoura and Llewellyn [24]. Acetonic extracts were filtered (porosity: 0.45 μm) and diluted volume-to-volume in 1 M ammonium acetate. The HPLC protocol followed Mantoura and Llewellyn [24] adapted by Méléder et al. [7]. Sample volumes of 100 μL were injected and pigments were detected by photodiode array and characterized by their absorption spectra between 400 and 800 nm and their retention time [25, 26]. Quantification was carried out at 440 nm using previously established pigment calibration with purified standards following Brotas and Plante-Cuny [27]. For some pigments, the contents of their derivatives were summed: Chl *a* and its allomers = Chl *a*; chlorophyll *c*₁ and *c*₂ = Chl *c*; fucoxanthin and fucoxanthin by-products = Fuco. Data were presented as cellular pigment contents ($\mu\text{g}\cdot\text{cell}^{-1}$), weight to weight pigment ratios ($\text{g}\cdot\text{g}^{-1}$) and pigment content per unit area ($\text{mg}\cdot\text{m}^{-2}$).

2.4 Absorption cross-section estimation

The MPBOM quantitative radiative transfer model was used to estimate a^* using the programming language IDL (Interactive Data Language) on ENVI 4.3 software. This model requires the input of biofilm reflectance spectra and the background reflectance spectra (i.e. wet filters without cells). One of the model outputs model is the absorption coefficient (α). This is related to the weighted concentration (C_i) and to the specific extinction coefficient (ε_i) at any wavelength of the m pigments present in the cells (Eq. 3):

$$\alpha = \sum_i^m \varepsilon_i C_i \quad (3)$$

α is dimensionless but when divided by the amount of Chl *a* (in $\text{mg}\cdot\text{m}^{-2}$) it corresponds to the absorption cross-section, a^* ($\text{m}^2\cdot\text{mg Chl}a^{-1}$). $\overline{a^*}$ corresponds to the averaged optical cross-section for the PAR domain, whereas $\overline{a^*}_{red}$ corresponds to the average for the domain of the red absorption of Chl *a* between 670 and 685 nm. For comparison between light treatments, $\Delta\overline{a^*}$ was calculated for each species. This measures the variation in absorption properties between LL and HL conditions and is associated with differences in Chl *a* ($\Delta\text{Chl } a$) cellular content.

All statistical analyses and graphical results were carried out in R statistical computing environment (2.6.1 [28]). Data normality was tested with the Shapiro-Wilk test while the effects of light acclimation (HL vs. LL) were tested using Student's t-test.

3 Results

3.1 Pigment analysis by HPLC

3.1.1 Pigment composition

In total, 14 liposoluble pigments (3 chlorophylls and 11 carotenoids) were detected and identified in this study (Table 1). *D. tertiolecta* showed the most diverse pigment composition (8 pigments), followed by *E. paludosa* and *S. platensis* (6 pigments) while only 3 pigments were detected in *P. cruentum*. Chl *a* and β -carotene (β -car) were detected in all organisms, while other pigments were species-specific: diadinoxanthin (DD), diatoxanthin (DT), fucoxanthin (Fuco) and Chl *c* were present in *E. paludosa* only; neoxanthin (NX), antheraxanthin (AX), lutein (Lut) and Chl *b* only in *D. tertiolecta*; oscillaxanthin (Osc) and myxoxanthophyll (Myx) only in *S. platensis*. In

contrast, *P. cruentum* did not exhibit any specific liposoluble pigments showing only Chl *a*, β -car and zeaxanthin (ZX).

3.1.2 Pigment changes in response to light

Light did not seem to have a large effect on the qualitative pigment composition. Only few pigments were detected exclusively in one of the two light conditions: violaxanthin (VX) in HL-acclimated *S. platensis* cells, Chl *b* in LL-acclimated *D. tertiolecta* cells and β -car in LL-acclimated cells for all the species except *P. cruentum*. It was assumed that the apparent lack of Chl *b* and β -car detection in HL-acclimated species was due to concentrations below the PDA detection threshold, rather than a true absence of pigments in the cells.

The cellular Chl *a* content for each species significantly decreased (t-test, $p < 0.05$) from LL to HL conditions, illustrating the photoacclimation effect (Table 2). Considering the other detected pigments, most of their cellular content followed the same trend with a significant decrease (t-test, $p < 0.05$) from LL to HL condition, with the exception of DT in *E. paludosa*, VX in *S. platensis* and ZX in *P. cruentum* which significantly increased ($p < 0.05$), AX in *D. tertiolecta* and DD in *E. paludosa* which remained stable ($p > 0.05$) (results not shown). The resulting pigment/Chl *a* ratios for each species at each irradiance are presented in Table 3. Notably, only Fuco:Chl *a* and Chl *c*:Chl *a* ratios (in *E. paludosa*) were not affected by the growth irradiance.

3.1.3 Pigment content on filters

The amount of Chl *a* collected on the different filters varied from $0.56 \text{ mg.m}^{-2} \pm 0.09$ on *S. platensis* HL-filters to $18.98 \text{ mg.m}^{-2} \pm 1.26$ on *E. paludosa* LL-filters (Figure 1). The amount of cells obtained by filtration on LL-acclimated filters was lower or similar

to HL filters (Table 2), thus the significant increase of pigment contents on LL-filters could only be the result of the increase in pigment cellular contents as a consequence of the light treatments. Only for *D. tertiolecta* and *P. cruentum* few pigments remained stable on filters (t-test, $p = 0.5$, figure 1).

For *D. tertiolecta* filters, Chl *b*, VX, and β -car followed the same trend as Chl *a*, and the opposite of AX (Figure 1a). Moreover, Lut, ZX and NX contents were comparable in HL and LL filters ($p = 0.48$, 0.23 or 0.25 , respectively). In *E. paludosa*, the content of Fuco, Chl*c* and β -car was higher in LL-filters, while it was similar for DD ($p = 0.6$) and significantly higher in HL-filters for DT (Figure 1b). *S. platensis* showed the largest Chl *a* decrease (up to 5.5 times) from LL to HL-filters and all the other pigments followed the same trend ($p \leq 0.05$) (Figure 1c). Finally, pigment contents of HL and LL-filters of *P. cruentum* were not significantly different ($p > 0.1$).

3.2 Radiometric analysis

Standardized reflectance spectra showed a similar shape whatever the species and the growth light conditions (Figure 2). A large absorption band in the blue domain, from 400 to 550/600 nm, was clearly visible followed by a transmittance peak and a second absorption band in the red domain around 675 nm. The transmittance peak varied in shape and was notched by smaller absorption bands: three in *D.tertiolecta* (Figure 2a), two in *E. paludosa* (Figure 2b), and one in *S. platensis* and *P. cruentum* (Figures 2c and d). Overall, reflectance values were lower in LL-acclimated cells indicating a higher absorbance. Almost no marked differences between species or light acclimation conditions could be inferred from the reflectance spectra, with the exception of the more pronounced 480 nm absorption band in LL-acclimated *D.tertiolecta* cells, and the specific 636 nm absorption band in *E. paludosa* (Figure 2).

The analysis of second derivative spectra improved the comparison between samples because each absorption peak could be assigned to an absorption band (Figure 3).

Some of the second derivative peaks were detected in all species, whatever the growth irradiance, while others were species- and growth irradiance-specific. Ubiquitous peaks were (Figure 3, downward arrows): ~ 416 nm; ~ 440 nm; ~ 462 (except LL-acclimated *D. tertiolecta* and *S. platensis* cells); ~ 546 nm and ~ 586 nm; ~ 625 nm (with a high variability from 615 to 631 nm), with two shoulders detected at 615 and 631 nm for LL-acclimated *P. cruentum* cells. An absorption double peak was detected in the red domain (~ 675 nm) in all species but *E. paludosa* (Figure 3). This double peak appeared in both photoacclimation conditions and was separated by a dip centered at 675 nm. The first peak (671 nm) was always smaller than the second one (683 nm) (Figure 3). Specific peaks (Figure 3, upward arrows) were: 1) ~ 489 nm in *D. tertiolecta* cells, which was smaller in LL-acclimated cells (Figure 3a) and was also present in HL-acclimated *S. platensis* cells (Figure 3c) and *P. cruentum* cells whatever the photoacclimation conditions (Figure 3d); 2) ~ 650 nm in *D. tertiolecta* cells (Figure 3a) and in HL-acclimated *S. platensis* cells (Figure 3c); 3) ~ 497 nm in LL-acclimated cells of *E. paludosa* (Figure 3b); 4) ~ 636 nm in *E. paludosa* cells (LL and HL) (Figure 3b); 5) ~ 524 nm in *S. platensis* (Figure 3c) and 6) ~ 570 nm in *P. cruentum* (LL and HL), which was smaller in LL-acclimated cells (Figure 3d).

3.3 Light absorption properties

Average absorption cross-section values through the PAR domain ($\overline{a^*}$) estimated using the MPBOM varied from 0.007 (*E. paludosa*) to 0.047 (*P. cruentum*) m².mg Chl a⁻¹ in LL-acclimated cells and from 0.024 (*E. paludosa*) to 0.117 (*S. platensis*) m².mg

Chl a^{-1} in HL-acclimated cells (Table 4). In the Chl a red absorption domain, i.e. between 670 and 685 nm, $\overline{a^*_{red}}$ vary from 0.008 (*E. paludosa*) to 0.063 (*S. platensis*) $m^2.mg$ Chl a^{-1} in LL-acclimated cells and from 0.027 (*E. paludosa*) to 0.106 (*S. platensis*) $m^2.mg$ Chl a^{-1} in HL-acclimated cells. Globally, values were greater in HL-acclimated cells, and $\overline{a^*_{red}}$ values were greater than $\overline{a^*}$ values, except *S. platensis* (Table 4). *E. paludosa* and *P. cruentum* showed a similar decrease in cellular Chl a content (-67 %, Table 4) but contrasting results regarding $\overline{a^*}$ and $\overline{a^*_{red}}$. In *E. paludosa* this decrease was accompanied by a 3-fold increase in $\overline{a^*}$ and $\overline{a^*_{red}}$ (reaching 226 %, Table 4) while in *P. cruentum* there was no significant changes in $\overline{a^*}$ and $\overline{a^*_{red}}$ regardless of the photoacclimation state.

4. Discussion

In many ecophysiological studies dealing with primary production and photosynthetic organisms with different pigment contents and photoacclimation strategies, the determination of the quantity of light absorbed and used to drive photosynthesis is a real challenge to estimate the photosynthetic efficiency, especially for field studies. The method herein described offers a simple way to determine a key factor in photobiology and photosynthetic studies, the optical absorption cross-section, a^* [2], by combining HPLC and reflectance data.

4.1 Advantages of using reflectance and second derivative spectra to assess the absorption properties of photosynthetic micro-organisms

In vivo measurement of the absorption properties of photosynthetic micro-organisms using spectroradiometry on cells settled as a biofilm is an efficient alternative to more classic approaches using spectrophotometry which suffer from light scattering and/or the need for β -correction [3, 5, 11-13]. Kazemipour et al. [14] showed that pigment reflectance mirrors well its absorbance when measurements are performed on a thin homogenous compact layer of micro-algal cells limiting distance between cells deposited on filters [14], but not thick enough to avoid saturation of reflectance measurements [7]. The biofilms used in the current study fitted these requirements because if there had been any scattering between cells, the model MPBOM would have not fitted the data. Additionally, the biomass was low enough to limit self-shading because in our experimental biofilms this is only observed at biomasses higher than 100 mg.Chl a.m⁻² [7].

Since light scattering and inherent reflectance are minimized, the reflectance measured this way corresponds to a double transmittance through the layer of cells. The incident light (i.e. PAR) is depleted by the absorbing fraction during the first transmission, reflected by the background, then transmitted again with a second depletion (see [14] for details). When the background reflectance is removed by standardization, the remaining reflectance signal contains only the cell absorption information.

To date, reflectance or absorbance spectra have been used to identify only a few specific pigment absorption features (e.g. the 650 nm band characteristic of the Chlorophyta or the 636 nm band of the Bacillariophyta). One of the major difficulties of assessing the *in vivo* pigment absorption properties is the possibility of existing multiple absorption bands for each pigment. Several factors may influence the pigment absorption properties in cells, e.g. the light scattering by the algal cell (or any

particle), the optical properties of each pigment, the spatial arrangement of pigments in LHCs and the spatial arrangement of cells within a sample [3]. All these factors can affect each other. Derivation of reflectance spectra helps to remove unwanted signals and improve the separation of overlapping pigment absorption features [29]. It can thus facilitate the identification of micro-organism absorption bands and their corresponding pigments.

In this study, the HPLC analysis of the 4 species revealed pigment signatures which are in agreement with current literature, for example regarding the clear effect of light observed on pigment ratios ([30] and references therein). Overall, most of the pigments detected by HPLC could be linked to specific derivative peaks. However, when a pigment/Chl *a* ratio was lower than 0.1, the pigment absorption was not strong enough to generate any derivative peak. This is why, for instance, VX absorption showed no second derivative peaks, although it is well-known to absorb at 510 nm *in vivo* [31].

4.2 Identification of chlorophyll absorption features

Second derivative reflectance peaks detected in each species and light condition were assigned to specific pigment absorption features on the basis of *in vitro* absorption bands and by the pigment changes induced by light acclimation (Table 1). The peaks most easily identified were those due to the ubiquitous pigment Chl *a*. In organic solvents, Chl *a* absorption bands are commonly detectable at 417, 431, 618 and 665 nm [32] and could be easily linked to the *in vivo* bands detected in all species at 416, 440, and 625 nm and to the often-observed second derivative double peak in the red domain (671 and 683 nm). This observation is similar to what was

observed by Hoepffner and Sathyendranath [5] using *in vivo* phytoplankton absorption spectra decomposed by a Gaussian function. These bands correspond to the major absorption bands according to the four-orbital model of Gouterman [33], i.e. the blue absorption bands (B-bands or Soret bands) and the red absorption bands (Q-bands) [2, 34]. The first two bands (416 and 440 nm) correspond to the Soret bands (B_y and B_x , respectively) whereas the last two bands (625 and around 675 nm) correspond to the Q-bands (Q_x and Q_y , respectively). The Q_y Chl *a* is responsible for all photochemical reactions due to its de-excitation to the ground state [2]. It is also in the Q_y transition band that a double peak was always observed at 671 and 683 nm, except in the diatom *E. paludosa*. However, this double absorption band was described by Jesus et al [19] in the same species, and by Serôdio et al [35] in natural biofilms. The double Q_y absorption peak is rarely mentioned in the literature and its origin remains unclear. Gulyayev and Litvin [18] were the first to report it at the same wavelengths as in the present study, using a variety of photosynthetic organisms, but they did not explain its origin. The same is true for more recent reports (e.g [2]) that also provided no explanation. To our knowledge, only Serôdio et al. [35] have suggested that the two peaks are the result of Chl *a* fluorescence emission centered at 683 nm. The chlorophyll fluorescence emitted at this wavelength would affect the radiometric data by increasing the reflectance at this wavelength and decreasing the absorption peak in the second derivative spectra [35]. However, it is possible that fluorescence emission is not the single factor affecting the dip between the two peaks. Another possible explanation for the double peak are absorption variations caused by photosystem (PS) stoichiometry. This is supported by previous observations that variations in PS stoichiometry (PSI and PSII) may cause a shift in the absorption from 674 to 678nm [5, 13]. In fact, because the PSI Chl *a* core

absorbs at 683 nm and the core of PSII Chl *a* at 670 nm, a change in the relative amount of the PSI against the PSII is likely to induce this shift [5]. The consequence in second derivative spectra should be the double peak, attributed to the two photosystem cores. The PSI:PSII ratio is often different from 1:1 and varies with species and with environmental light conditions [5, 36-38]. This mechanism is a long-term photoacclimation process that requires hours to days to occur. It allows the excitation energy within the photosynthetic machinery to be regulated by changing the relative amount of the two photosystems [39]. For example, in this study, the Chlorophyta species showed the largest difference between light conditions with the second derivative peak at 671 nm increasing more than the 683 nm in the HL treatment. This supports the hypothesis that HL-acclimated *D. tertiolecta* cells decreased their PSI:PSII ratio as proposed by Walters et al. [38]. The Rhodophyta and Cyanophyta species showed similar height for the two peaks in both light conditions, which could be explained by a stable PSI:PSII ratio. This might be cause by a parallel decrease in both photosystems when the cells are HL-acclimated as reported previously for Rhodophyta [40]. Although the assessment of PS stoichiometry by radiometry looks very promising further studies are still required to test our hypothesis, using for example, reflectance analyses of cells with controlled amount of PSI and PSII

Three ubiquitous peaks (466, 546 and 586 nm) may also be assigned to Chl *a* as they are minor absorption bands observed *in vitro* in organic solvent, with a shift [41].

This assumption is based on the fact that Chl *a* was the only pigment found in the four species. However, some uncertainty remains in the diatom species since Fuco is known to exhibit a broad absorption band around 550 nm *in vivo* [42]. Thus, a

pigment mixture including Chl *a* could also explain the presence of these minor ubiquitous bands.

The other chlorophylls (*b* and *c*) also show a Q_y band (636 nm for Chl *c* and 650 nm for Chl *b*) without any double peak feature. These Q_y absorption bands are in accordance with previous findings [5, 7, 43]. The other major absorption bands (Soret bands and Q_x) were not observed, possibly because of their small height and/or because of their overlap with other absorption peaks.

4.3 Identification of carotenoid absorption features

Carotenoids are structurally and functionally the most diverse pigment group. They cover part of the 'green gap' with absorption bands from 450 to 550 nm [44]. Several carotenoids are involved in photoacclimation and photoregulation processes, e.g. xanthophyll cycles and structural adjustment of the LHCs [39]. Xanthophylls are important in micro-algal photoprotective processes [45], which explains the increase in the xanthophyll:Chl *a* ratio in the Chlorophyta (VX, AX and ZX pool) and the diatom (DD and DT pool) under HL. This led to a higher absorption in the 489-500 nm region. In the Chlorophyta, the absorption increase at 489 nm was caused by the presence of the xanthophylls ZX and Lut, with absorption bands at 485 nm [39] and 494 nm [9, 30], respectively. The 489 nm peak was also observed in the Cyanophyta and the Rhodophyta species, both containing ZX. In the diatom species, the absorbance was centered at 497 nm. This was probably due to the joint absorption of DD and DT at 487 and 508 nm, respectively [10, 16, 46]. The case of the xanthophyll pigments illustrates very well the difficulty in deconvoluting the individual contribution of pigments with very similar reflectance spectrum shapes. Nevertheless, some authors have demonstrated the possibility of obtaining very detailed information from

spectral data. For instance, Jesus et al. [16] observed a shift from 487 to 508 nm associated with the operation of the xanthophyll cycle which corresponds to the HL enzymatic conversion of DD to DT, while Ruban et al. [10] and Lavaud and Lepetit [47] have shown that a thin spectral band around 520 nm is related to the DT molecules which effectively involved in the photoprotective dissipation of excess energy (the so-called NPQ, i.e. the non photochemical fluorescence quenching).

In the Cyanophyta species, the absorption features between 450-550 nm showed the largest differences between the two light acclimation treatments. The increase in the 489 nm absorption band from LL to HL was probably caused by the increase in ZX. The specific absorption band at 524 nm increased from LL to HL, which could be caused by increases in either Myx or Osc. However, Myx is the most likely source of this peak because the amount of Osc was 5 times lower than Myx and was probably too low to be retrieved from the spectral data in a reliable way.

4.4 Identification of phycobiliproteins (PBPs) absorption features

Although PBPs were not quantified in the current study, some absorption bands could be linked to this pigment group. The Cyanophyta species showed a series of absorption bands from ~585 to ~630 nm, which could correspond to the absorption domains of the blue PBP phycocyanins (610-620 nm) and allophycocyanins (650-655 nm) [48]. Allophycocyanin peaks appeared to show an unexpected increase in HL-acclimated cells of the Cyanophyta species, which is not consistent with their known light-harvesting function [49]. However, under high light condition a phosphorylation cascade is induced in cells activating specific polypeptides that degrade PBPs [50], which is accompanied by a Chl *a* decrease. A greater allophycocyanin/Chl *a* ratio illustrated by a higher absorption peak at 650 nm

suggests that the decrease of the PSI and PSII (containing Chl *a*) was faster than the decrease of PBPs, whereas the expected trend is a decrease of the ratio with light [40]. The same unexpected trend was observed with phycocyanin in the Rhodophyta species, where HL-acclimated cells showed higher absorption peaks around 570 nm corresponding to the absorption by phycoerythrin [48].

4.5 Estimation of optical absorption cross-section (a^*)

One goal of this study was to demonstrate the possibility of assessing a^* from reflectance data using the MPBOM [14] regardless of the taxonomic algal group considered. This parameter is required for the calculation of the absolute photosynthesis quantum yield which is essential to estimate primary productivity from incident PAR. Also, it must be estimated *in vivo* because a^* is highly dependent on cell size and pigment composition as well as on plastid size and number, which drive the pigment-package effects, i.e. self-shading, light scattering, etc. [2]. By settling the micro-algal cells on a filter, the light scattering effects observed in a cell suspension are suppressed and thus allowing to use reflectance to estimate a^* *in vivo*. This approach avoids the use of complex apparatus, as an integrating sphere. Values of $\overline{a^*}$ and $\overline{a^*_{red}}$, estimated by the MPBOM were in accordance with published values, as well as changes observed with cells grown under LL or HL increase has also been reported before [13]. The efficiency of light absorption in HL-acclimated cells is regulated on a long-term scale (i.e. several days, depending on the growth rate) by a decrease in the number of thylakoids per plastid generating a decrease in the plastid volume and/or by a decrease in the number of plastids per cell (in species with many small plastids), and also by changes in the cellular pigment content. Both processes lead to a decrease in the “package effect” [2]. Often, these phenomena occur

together with a reduction in LHCs size, depending on the light conditions [2]. The diatom species showed the highest $\overline{a^*}$ increase with light, suggesting that this group might be the most efficient to acclimate to changing light conditions, possibly due to its strong potential for the regulation of photosynthesis in fluctuating light conditions [45, 51]. In contrast, the Rhodophyta species showed a stability in a^* values instead of the expected increase due to package effect (rearrangement of plastids, thylakoids and LHCs). In spite of the photoacclimation status (i.e. decrease in Chl *a* cellular content from LL to HL), the cells showed the same absorption cross-section values and these results suggest that there was no rearrangement of plastids, thylakoids and/or LHCs for this taxonomical group. This can be explained by a specific photoacclimation strategy described for *P. cruentum* [40] whereby Chl *a* and phycobilisome content per thylakoid area remain relatively constant even though the overall pigment content decreases in HL-acclimated cells.

Another relevant result observed in the Chlorophyta and the Cyanophyta species was the a^* increase from LL to HL, which was highest when the average was calculated over the PAR domain in comparison to what was measured in Chl *a* absorption red domain. Between 670 and 680 nm, it is known that only Chl *a* absorbs the light energy, whereas in the rest of the PAR spectrum, carotenoids also absorb, including the photoprotective ones [13] as for instance violaxanthin. Thus, the coupled increase in photoprotective pigments in HL photoacclimated cells and the Chl *a* decrease, could explain the bigger $\overline{a^*}$ values in comparison to the $\overline{a^*_{red}}$ values. Consequentially, the a^* increase seems to be not only due to the rearrangement of the package effect, but also to the increase in photoprotective pigments, although they could not participate in the energy transfer to the cores Chl *a* [13]. Indeed, some of these pigments are only involved in the xanthophyll cycle, as AX, ZX and DT.

It should also be noted that \overline{a}_{red}^* did not show significant difference between LL and HL conditions in the Chlorophyta species in contrast to \overline{a}^* . This implies that the \overline{a}^* increase was only caused by the carotenoid increase and in this case the package effect affected only \overline{a}_{red}^* , reducing its decrease in spite of the cellular Chl *a* content. This result was in accordance with those found by Johnsen and Sakshaug [13] that observed that Chlorophyta are also the species with the lowest absorption-cross section coefficients.

5. Conclusion and future prospects

The non-invasive radiometric methodology proposed in this study was shown to be a good alternative to the already existing techniques for the assessment of the pigment absorption properties of photosynthetic micro-organisms. Namely, *in vivo* pigment absorption bands could be identified from second derivative reflectance spectra in a reliable way. More importantly, by coupling this approach to the MPBOM model, we demonstrated the feasibility of measuring the absorption cross-section (a^*) of photosynthetic microorganisms *in vivo*, which is a key parameter for the estimation of the absolute photosynthetic efficiency and the primary productivity. In agreement with previous works, the pattern of variation of a^* observed in this study suggests that these photosynthetic microorganisms display a high plasticity in response to light changes, and that the differences in responses between taxonomic groups are large [52]. Last but not least, this non-invasive methodology will facilitate the investigation of pigment absorption properties of microphytobenthos biofilms *in vivo* and *in situ*. An example of an original application could be the mapping of microphytobenthos light absorption efficiency using remote sensing imagery. This would strongly help assessing coastal primary production at the habitat/ecosystem scale, such as intertidal mudflats.

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Figure legends

Figure 1. Pigment concentration expressed in mg.m^{-2} in the four species acclimated at the two light (PAR) intensities, LL (black) and HL (gray). For each pigment (mean \pm 95 % CI; $n = 3$), horizontal bar indicates no significant differences between HL and LL conditions (t test $p \geq 0.05$).

Figure 2. Spectra of standardized reflectance (ρ_{925}) measured for the four species acclimated at the two light (PAR) intensities, LL (black) and HL (gray). Arrows indicate absorption bands notching the transmittance peak of the 'green gap'.

Figure 3. Spectra of second derivative ($\delta\delta$) calculated for the four species acclimated at the two light (PAR) intensities, LL (black) and HL (gray). Downward arrows indicate ubiquitous absorption peaks while upward arrows indicate specific peaks.

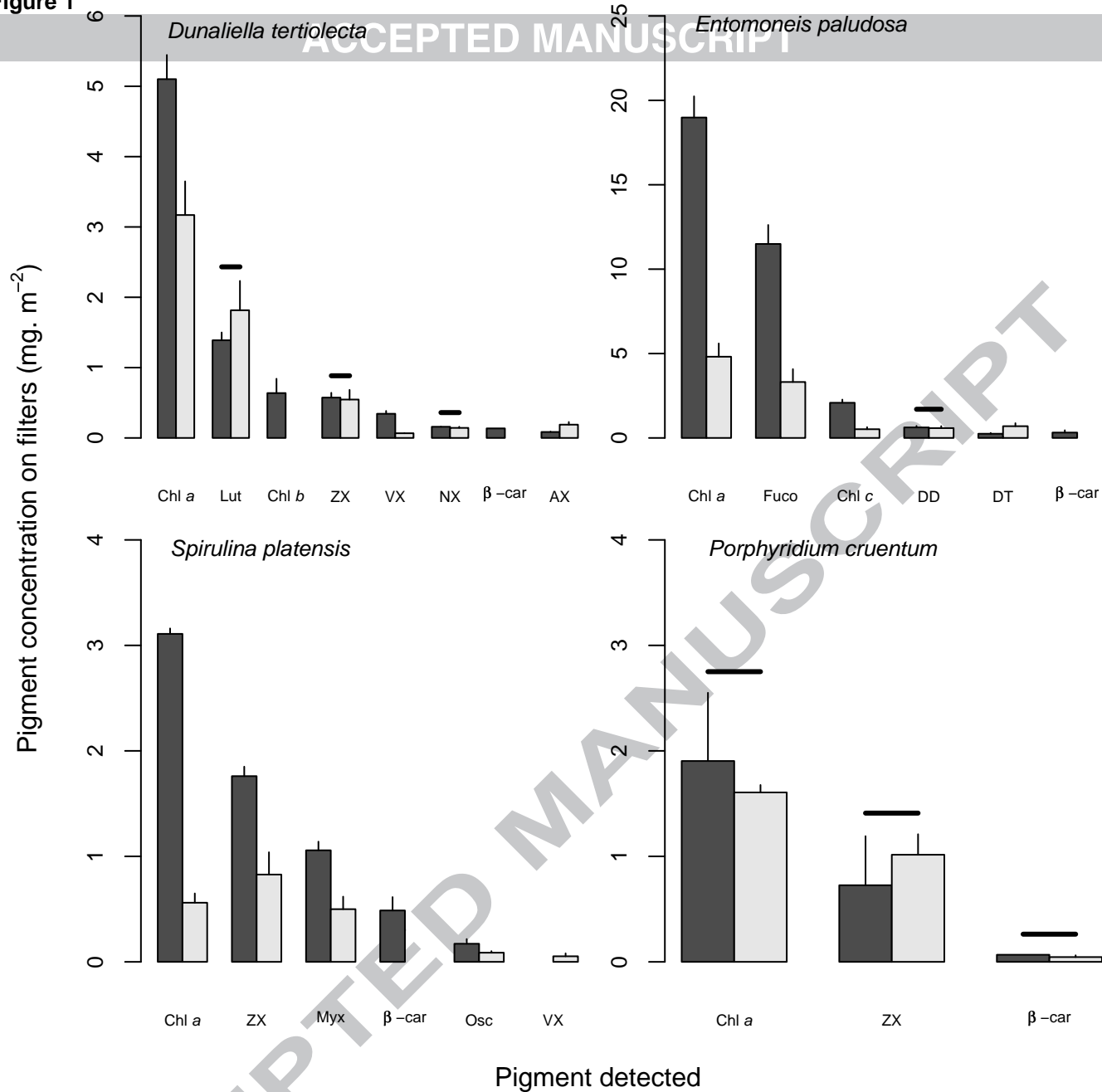
Figure 1

Figure 2

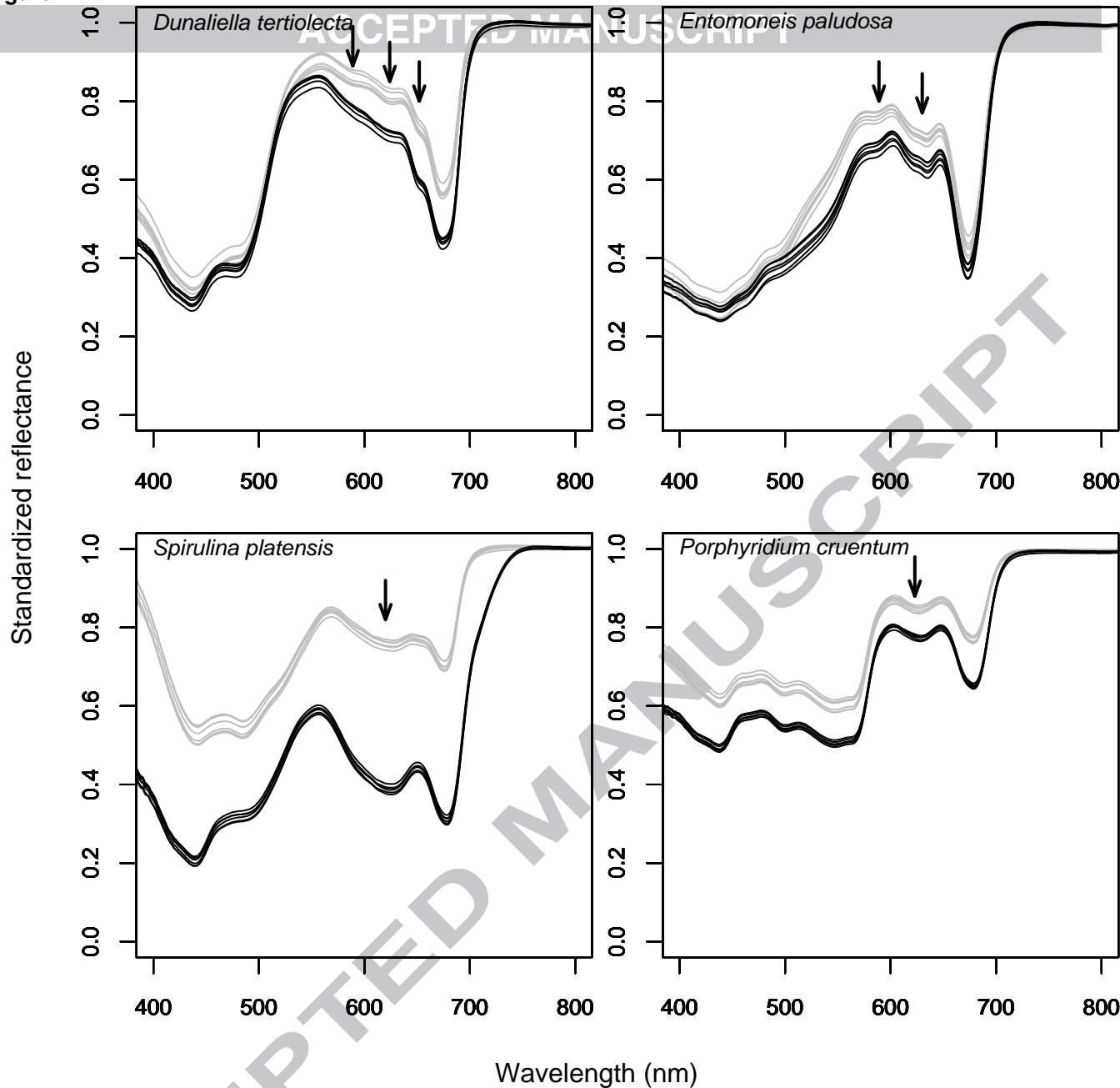
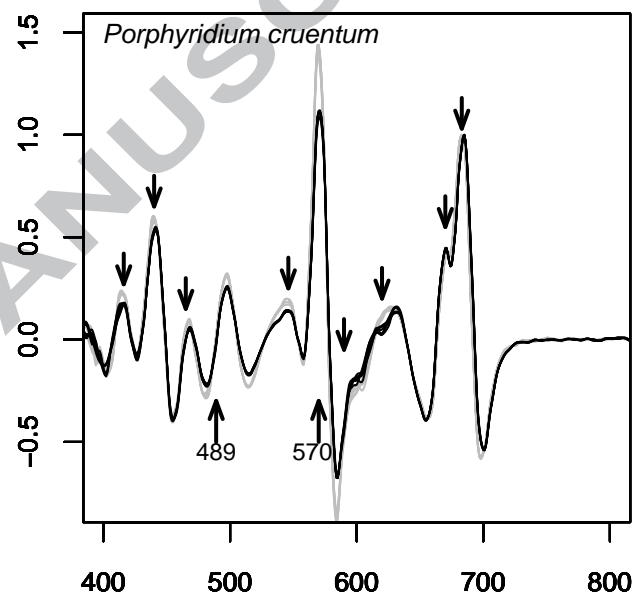
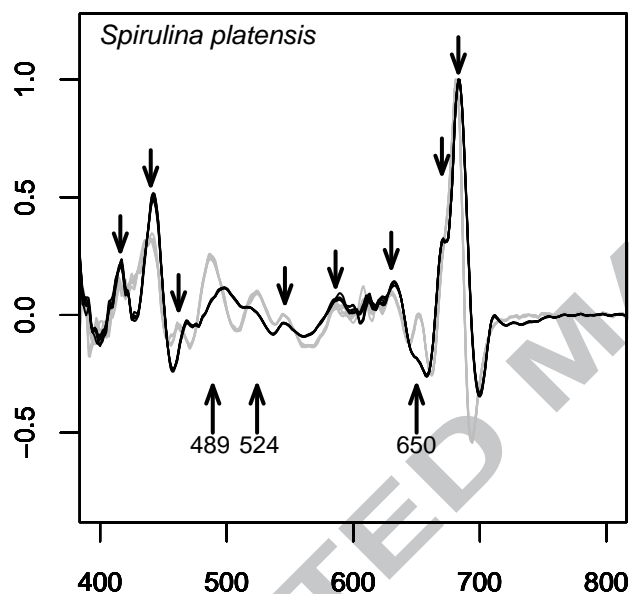
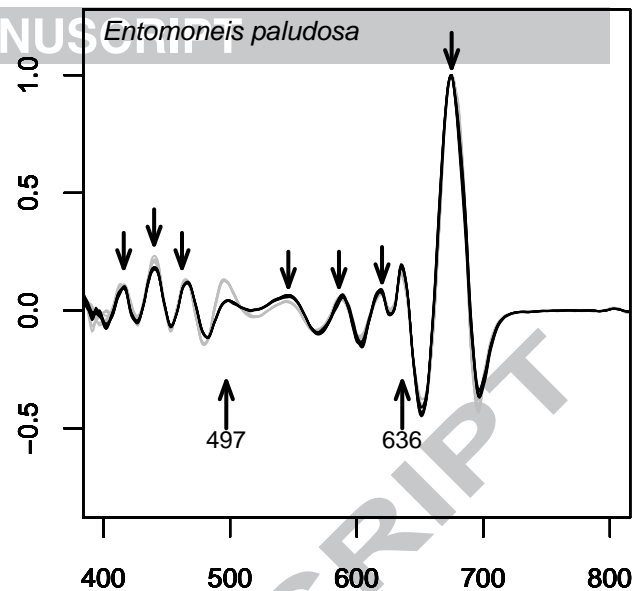
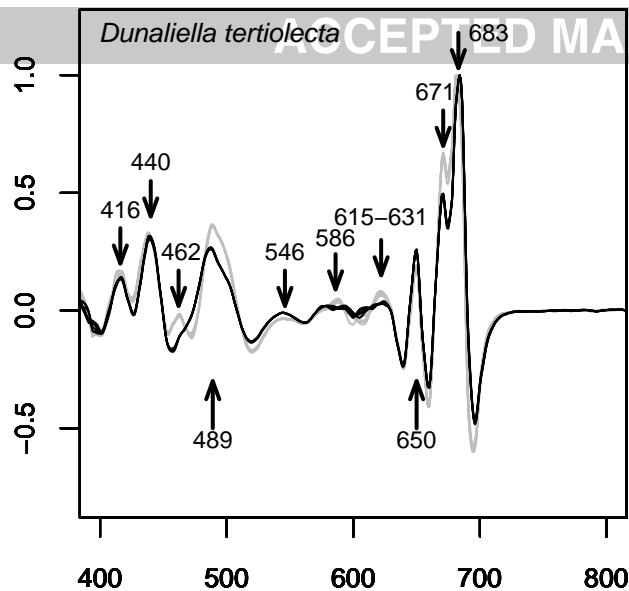


Figure 3

Standardized Second Derivative



Wavelength (nm)

Table 1. Absorbance maxima for all pigments detected in the four strains analyzed by HPLC (*in vitro*) and by radiometry (*in vivo*). Pigments are ordered following their HPLC elution time from the earliest to the latest. n.d.: *in vivo* absorbances not detected by radiometry. References for *in vitro* absorbances: Jeffrey et al. [26], except (*) in Brotas and Plante-Cuny, [25]. *In vivo* absorbance are from this study.

Pigment	Maxima <i>in vitro</i>			Maxima <i>in vivo</i>		Sources
Chlorophyll <i>c</i> (Chl <i>c</i>)	442	573	630	636		<i>E. paludosa</i>
Fucoxanthin (Fuco)	451 (465)			550		<i>E. paludosa</i>
Oscillaxanthin (Osc)*	(469)	498	529	n.d.		<i>S. platensis</i>
Neoxanthin (NX)	(414)	437	466	n.d.		<i>D. tertiolecta</i>
Violaxanthin (VX)	(417)	440	469	n.d.		<i>D. tertiolecta</i> , <i>S. platensis</i>
Myxoxanthophyll (Myx)*	(451)	475	508	524		<i>S. platensis</i>
Diadinoxanthin (DD)	(420)	447	477	DD+DT: 496		<i>E. paludosa</i>
Antheraxanthin (AX)	(421)	444	471	n.d.		<i>D. tertiolecta</i>
Diatoxanthin (DT)		453	481	DD+DT: 496		<i>E. paludosa</i>
Lutein (Lut)	(421)	446	474	500		<i>D. tertiolecta</i>
Zeaxanthin (ZX)		452	478	489		<i>D. tertiolecta</i> , <i>S. platensis</i> (HL), <i>P. cruentum</i>
Chlorophyll <i>b</i> (Chl <i>b</i>)		458	596	646	650	<i>D. tertiolecta</i>
Chlorophyll <i>a</i> (Chl <i>a</i>)	431		(581)	618	625	All 4 strains
β -Carotene (β -car)	425	453	481	466	499	All 4 strains

Table 2. Cellular content of Chl *a* in the four studied species photoacclimated at the two PAR intensities, HL and LL. For each value (mean \pm 95 % CI; *n* = 3), differences between LL and HL are significant (t test $p \leq 0.05$).

Strain	Light acclimation	Cellular content of Chl <i>a</i> ($\mu\text{g} \cdot 10^{-6}$ cell)	Amount of cells on filters
<i>D. tertiolecta</i>	LL	4.37 ± 1.08	$2.4 \cdot 10^6 \pm 2.0 \cdot 10^5$
	HL	1.02 ± 0.15	$5.5 \cdot 10^6 \pm 2.7 \cdot 10^5$
<i>E. paludosa</i>	LL	11.45 ± 0.76	$2.9 \cdot 10^6 \pm 1.5 \cdot 10^5$
	HL	3.78 ± 0.62	$2.3 \cdot 10^6 \pm 2.2 \cdot 10^5$
<i>S. platensis</i>	LL	0.69 ± 0.26	$1.0 \cdot 10^7 \pm 3.1 \cdot 10^5$
	HL	0.09 ± 0.01	$1.2 \cdot 10^7 \pm 3.5 \cdot 10^5$
<i>P. cruentum</i>	LL	0.65 ± 0.13	$4.0 \cdot 10^6 \pm 3.2 \cdot 10^5$
	HL	0.21 ± 0.01	$1.3 \cdot 10^7 \pm 1.1 \cdot 10^6$

Table 3. Pigment content expressed in ratio against Chl *a* in the four studied species photoacclimated at the two PAR intensities, HL and LL. For each ratio (mean \pm 95 % CI; *n* = 3), differences between LL and HL are significant (t test $p \leq 0.05$), except (n.s.).

Ratio \pm std	<i>D. tertiolecta</i> LL	<i>D. tertiolecta</i> HL
Lut / Chl <i>a</i>	0.27 \pm 0.01	0.57 \pm 0.05
Chl <i>b</i> / Chl <i>a</i>	0.13 \pm 0.03	n.d.
ZX / Chl <i>a</i>	0.11 \pm 0.005	0.17 \pm 0.005
VX / Chl <i>a</i>	0.07 \pm 0.003	0.02 \pm 0.003
NX / Chl <i>a</i>	0.03 \pm 0.001	0.05 \pm 0.003
β -car / Chl <i>a</i>	0.08	n.d.
AX / Chl <i>a</i>	0.02 \pm 0.003	0.06 \pm 0.003
	<i>E. paludosa</i> LL	<i>E. paludosa</i> HL
Fuco / Chl <i>a</i> (n.s.)	0.61 \pm 0.03	0.69 \pm 0.07
Chl <i>c</i> / Chl <i>a</i> (n.s.)	0.11 \pm 0.007	0.11 \pm 0.01
DD / Chl <i>a</i>	0.03 \pm 0.002	0.12 \pm 0.01
DT / Chl <i>a</i>	0.01 \pm 0.002	0.14 \pm 0.02
β -car / Chl <i>a</i>	0.02 \pm 0.008	n.d.
	<i>S. platensis</i> LL	<i>S. platensis</i> HL
ZX / Chl <i>a</i>	0.57 \pm 0.02	1.46 \pm 0.15
Myx / Chl <i>a</i>	0.34 \pm 0.02	0.88 \pm 0.08
β -car / Chl <i>a</i>	0.16 \pm 0.04	n.d.
Osc / Chl <i>a</i>	0.06 \pm 0.01	0.16 \pm 0.04
VX / Chl <i>a</i>	n.d.	0.09 \pm 0.04
	<i>P. cruentum</i> LL	<i>P. cruentum</i> HL
ZX / Chl <i>a</i>	0.37 \pm 0.13	0.63 \pm 0.1
β -car / Chl <i>a</i>	0.03	0.07 \pm 0.02

Table 4. Absorption cross-section estimated from MPBOM for the four studied strains photoacclimated at the two PAR intensities, HL and LL. $\overline{a^*}$: absorption cross-section in $\text{m}^2.\text{mg Chl } a^{-1}$, averaged on the PAR domain $\pm 95\%$ CI; $\overline{a^*_{red}}$: absorption cross-section in $\text{m}^2.\text{mg Chl } a^{-1}$, averaged between 670 and 685 nm; $\Delta\overline{a^*}$, $\Delta\overline{a^*_{red}}$ and $\Delta\text{Chl } a$: variation from LL to HL conditions in cells. For each strain, differences between LL and HL are significant (t test $p \leq 0.05$), except n.s ($\Delta\overline{a^*}$ and $\Delta\overline{a^*_{red}}$ were minimized to zero).

Strain	Light acclimation	$\overline{a^*}$ ($\text{m}^2.\text{mg Chl } a^{-1}$)	$\overline{a^*_{red}}$ ($\text{m}^2.\text{mg Chl } a^{-1}$)	$\Delta\overline{a^*}$	$\Delta\overline{a^*_{red}}$	$\Delta\text{Chl } a$
<i>D. tertiolecta</i>	LL	0.017 ± 0.003	0.023 ± 0.004	57 %	0 % n.s	-71 %
	HL	0.027 ± 0.003	0.030 ± 0.004			
<i>E. paludosa</i>	LL	0.007 ± 0.000	0.008 ± 0.000	226 %	225 %	-67 %
	HL	0.024 ± 0.005	0.027 ± 0.005			
<i>S. platensis</i>	LL	0.054 ± 0.002	0.063 ± 0.002	117 %	68 %	-88 %
	HL	0.117 ± 0.021	0.106 ± 0.017			
<i>P. cruentum</i>	LL	0.047 ± 0.013	0.040 ± 0.011	0 % (n.s.)	0 % (n.s.)	-67 %
	HL	0.035 ± 0.002	0.027 ± 0.001			

Highlights :

- *In vivo* estimation of the pigment absorption bands and the optical absorption cross-section, a^*
- Four photosynthetic micro-organisms cultivated at two PAR levels were analyzed
- A new methodology was proposed: spectroradiometry measuring reflectances.
- Some absorption bands were ubiquitous, others were taxonomically specific and/or photo-physiological dependent.
- A double peak feature at 671 and 683 nm, supposed to be caused by PSII and PSI, was detected